#5

## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/501.8/4
Source:	PUTIO
Date Processed by STIC:	9/6/05

## ENTERED



PCT

RAW SEQUENCE LISTING DATE: 09/06/2005
PATENT APPLICATION: US/10/501,814 TIME: 11:28:50

```
3 <110> APPLICANT: Evotec NeuroSciences GmbH
      5 <120> TITLE OF INVENTION: Diagnostic and therapeutic use of a voltage-gated ion
              channel for neurodegenerative diseases
      8 <130> FILE REFERENCE: P67564US1
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/501,814
C--> 11 <141> CURRENT FILING DATE: 2004-07-19
     13 <160> NUMBER OF SEQ ID NOS: 15
     15 <170> SOFTWARE: PatentIn Ver. 2.1
     17 <210> SEQ ID NO: 1
     18 <211> LENGTH: 272
     19 <212> TYPE: DNA
     20 <213> ORGANISM: Artificial Sequence
     22 <220> FEATURE:
     23 <223> OTHER INFORMATION: Description of Artificial Sequence: cDNA fragment
              of the human SCN2A gene
     26 <400> SEOUENCE: 1
     27 aattaaggtt ggaagaataa aaagcaagaa gctcttcctt gtttgctgca acctattgct 60
     28 taatgacatg aagaatgagg tettggtaga acaatttget teaetttace actgatatat 120
     29 ggcttcccat attagacttc tgaacagggg aaggaataag atacagcagc ataggcaaga 180
     30 taaacatgca gcagtgacag ettcaaacta taatggaacc aattacatca tattacctgt 240
     31 tggaagettg caaactatac ttactggggt ac
     34 <210> SEQ ID NO: 2
     35 <211> LENGTH: 8292
     36 <212> TYPE: DNA
     37 <213> ORGANISM: Artificial Sequence
     39 <220> FEATURE:
     40 <223> OTHER INFORMATION: Description of Artificial Sequence: cDNA of the
              human SCN2A gene
     41
     43 <400> SEQUENCE: 2
     44 cactttetta tqeaaqqaqe taaacaqtqa ttaaaqqaqe aqqatqaaaa qatqqeacaq 60
     45 tragtgrigg tarregragg arctgarage tirregritet tracraggga atceptiget 120
     46 gctattgaac aacgcattgc agaagagaaa gctaagagac ccaaacagga acgcaaggat 180
     47 gaggatgatg aaaatggccc aaagccaaac agtgacttgg aagcaggaaa atctcttcca 240
     48 tttatttatg gagacattcc tccagagatg gtgtcagtgc ccctggagga tctggacccc 300
     49 tactatatca ataagaaaac gtttatagta ttgaataaag ggaaagcaat ctctcgattc 360
     50 agtgccaccc ctgcccttta cattttaact cccttcaacc ctattagaaa attagctatt 420
     51 aagattttgg tacattettt atteaatatg eteattatgt geaegattet taceaactgt 480
     52 gtatttatga ccatgagtaa ccctccagac tggacaaaga atgtggagta tacctttaca 540
     53 ggaatttata cttttgaatc acttattaaa atacttgcaa ggggcttttg tttagaagat 600
     54 ttcacatttt tacgggatcc atggaattgg ttggatttca cagtcattac ttttgcatat 660
     55 gtgacagagt ttgtggacct gggcaatgtc tcagcgttga gaacattcag agttctccga 720
     56 gcattgaaaa caatttcagt cattccaggc ctgaagacca ttgtgggggc cctgatccag 780
     57 tcagtgaaga agctttctga tgtcatgatc ttgactgtgt tctgtctaag cgtgtttgcg 840
```

## **RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/501,814**DATE: 09/06/2005 TIME: 11:28:50

```
58 ctaataggat tgcagttgtt catgggcaac ctacgaaata aatgtttgca atggcctcca 900
59 gataattett eetttgaaat aaatateact teettettta acaatteatt ggatgggaat 960
60 ggtactactt tcaataggac agtgagcata tttaactggg atgaatatat tgaggataaa 1020
61 aqtcactttt attttttaqa qqqqcaaaat qatqctctqc tttqtqqcaa cagctcagat 1080
62 qcaqqccaqt gtcctgaagg atacatctgt gtgaaggctg gtagaaaccc caactatggc 1140
63 tacacqaqct ttgacacctt tagttgggcc tttttgtcct tatttcgtct catgactcaa 1200
64 qacttctqqq aaaaccttta tcaactgaca ctacgtgctg ctgggaaaac gtacatgata 1260
65 ttttttgtgc tggtcatttt cttgggctca ttctatctaa taaatttgat cttggctgtg 1320
66 gtggccatgg cctatgagga acagaatcag gccacattgg aagaggctga acagaaggaa 1380
67 qctqaatttc aqcaqatqct cqaacaqttq aaaaagcaac aaqaaqaagc tcaggcggca 1440
68 gctgcagccg catctgctga atcaagagac ttcagtggtg ctggtgggat aggagttttt 1500
69 tcagagagtt cttcagtagc atctaagttg agctccaaaa gtgaaaaaga gctgaaaaac 1560
70 agaagaaaga aaaagaaaca gaaagaacag tctggagaag aagagaaaaa tgacagagtc 1620
71 cgaaaatcgg aatctgaaga cagcataaga agaaaaggtt tccgtttttc cttggaagga 1680
72 agtaggetga catatgaaaa gagattttet tetecaeaee agteettaet gageateegt 1740
73 ggctcccttt tctctccaag acgcaacagt agggcgagcc ttttcagctt cagaggtcga 1800
75 aatqacaqcc qaaqaqactc tctgttcgtg ccgcacagac atggagaacg gcgccacagc 1920
76 aatgtcagcc aggccagccg tgcctccagg gtgctcccca tcctgcccat gaatgggaag 1980
77 atgcatageg etgtggaetg caatggtgtg gteteeetgg tegggggeee ttetaceete 2040
78 acatetgetg ggeageteet accagaggge acaactactg aaacagaaat aagaaagaga 2100
79 cggtccagtt cttatcatgt ttccatggat ttattggaag atcctacatc aaggcaaaga 2160
80 gcaatgagta tagccagtat tttgaccaac accatggaag aacttgaaga atccagacag 2220
81 aaatgcccac catgctggta taaatttgct aatatgtgtt tgatttggga ctgttgtaaa 2280
82 ccatqqttaa aqqtqaaaca ccttqtcaac ctqqttqtaa tqqacccatt tqttqacctg 2340
83 qccatcacca tctqcattqt cttaaataca ctcttcatqq ctatqqaqca ctatcccatg 2400
84 acggagcagt tcagcagtgt actgtctgtt ggaaacctgg tcttcacagg gatcttcaca 2460
85 gcagaaatgt ttctcaagat aattgccatg gatccatatt attactttca agaaggctgg 2520
86 aatatttttg atggttttat tgtgagcctt agtttaatgg aacttggttt ggcaaatgtg 2580
87 gaaggattgt cagttctccg atcattccgg ctgctccgag ttttcaagtt ggcaaaatct 2640
88 tggccaactc taaatatgct aattaagatc attggcaatt ctgtgggggc tctaggaaac 2700
89 ctcaccttgg tattggccat catcgtcttc atttttgctg tggtcggcat gcagctcttt 2760
90 ggtaagaget acaaagaatg tgtetgeaag atttecaatg attgtgaact cecaegetgg 2820
91 cacatgcatg actttttcca ctccttcctg atcgtgttcc gcgtgctgtg tggagagtgg 2880
92 atagagacca tgtgggactg tatggaggtc gctggccaaa ccatgtgcct tactgtcttc 2940
93 atgatggtca tggtgattgg aaatctagtg gttctgaacc tcttcttggc cttgcttttg 3000
94 agttccttca gttctgacaa tcttgctgcc actgatgatg ataacgaaat gaataatctc 3060
95 cagattgctg tgggaaggat gcagaaagga atcgattttg ttaaaagaaa aatacgtgaa 3120
96 tttattcaga aagcctttgt taggaagcag aaagctttag atgaaattaa accgcttgaa 3180
97 gatctaaata ataaaaaaga cagctgtatt tccaaccata ccaccataga aataggcaaa 3240
98 gacctcaatt atctcaaaga cggaaatgga actactagtg gcataggcag cagtgtagaa 3300
99 aaatatqtcq tqqatqaaaq tqattacatq tcatttataa acaaccctag cctcactgtg 3360
100 acaqtaccaa ttqctqttqq aqaatctqac tttqaaaatt taaatactga agaattcagc 3420
101 agcgagtcag atatggagga aagcaaagag aagctaaatg caactagttc atctgaaggc 3480
102 agcacggttg atattggagc tcccgccgag ggagaacagc ctgaggttga acctgaggaa 3540
103 tecettgaac etgaageetg ttttacagaa gaetgtgtac ggaagtteaa gtgttgteag 3600
104 ataagcataq aagaaggcaa agggaaactc tggtggaatt tgaggaaaac atgctataag 3660
105 ataqtqqaqc acaattqqtt cgaaaccttc attqtcttca tgattctgct gagcagtggg 3720
106 gctctggcct ttgaagatat atacattgag cagcgaaaaa ccattaagac catgttagaa 3780
```

**RAW SEQUENCE LISTING**PATENT APPLICATION: **US/10/501,814**DATE: 09/06/2005

TIME: 11:28:50

```
107 tatgctgaca aggttttcac ttacatattc attctggaaa tgctgctaaa gtgggttgca 3840
108 tatggttttc aagtgtattt taccaatgcc tggtgctggc tagacttcct gattgttgat 3900
109 gtctcactgg ttagcttaac tgcaaatgcc ttgggttact cagaacttgg tgccatcaaa 3960
110 teceteaqaa caetaaqage tetgaggeea etgagagett tgteeeggtt tgaaggaatg 4020
111 agggttgttg taaatgctct tttaggagcc attccatcta tcatgaatgt acttctggtt 4080
112 tqtctqatct tttqqctaat attcaqtatc atqqqaqtqa atctctttqc tqqcaaqttt 4140
113 taccattqta ttaattacac cactqqaqaq atqtttqatq taaqcqtqqt caacaactac 4200
114 agtgagtgca aagctctcat tgagagcaat caaactgcca ggtggaaaaa tgtgaaagta 4260
115 aactttgata acgtaggact tggatatctg tetetactte aagtageeac gtttaaggga 4320
116 tggatggata ttatgtatgc agctgttgat tcacgaaatg tagaattaca acccaagtat 4380
117 gaagacaacc tgtacatgta tctttatttt gtcatcttta ttatttttgg ttcattcttt 4440
118 accttgaatc ttttcattgg tgtcatcata gataacttca accaacagaa aaagaagttt 4500
119 ggaggtcaag acatttttat gacagaagaa cagaagaaat actacaatgc aatgaaaaaa 4560
120 ctgggttcaa agaaaccaca aaaacccata cctcgacctg ctaacaaatt ccaaggaatg 4620
121 gtctttgatt ttgtaaccaa acaagtcttt gatatcagca tcatgatcct catctgcctt 4680
122 aacatggtca ccatgatggt ggaaaccgat gaccagagtc aagaaatgac aaacattctg 4740
123 tactggatta atctggtgtt tattgttctg ttcactggag aatgtgtgct gaaactgatc 4800
124 tctcttcgtt actactattt cactattgga tggaatattt ttgattttgt ggtggtcatt 4860
125 ctctccattg taggaatgtt tctggctgaa ctgatagaaa agtattttgt gtcccctacc 4920
126 ctgttccgag tgatccgtct tgccaggatt ggccgaatcc tacgwctgat caaaggagca 4980
127 aaggggatee geacgetget etttgetttg atgatgteee tteetgegtt gtttaacate 5040
128 ggcctccttc ttttcctggt catgttcatc tacgccatct ttgggatgtc caattttgcc 5100
129 tatgttaaga gggaagttgg gatcgatgac atgttcaact ttgagacctt tggcaacagc 5160
130 atgatctgcc tgttccaaat tacaacctct gctggctggg atggattgct agcacctatt 5220
131 cttaatagtg gacctccaga ctgtgaccct gacaaagatc accctggaag ctcagttaaa 5280
132 ggagactgtg ggaacccatc tgttgggatt ttcttttttg tcagttacat catcatatcc 5340
133 ttcctggttg tggtgaacat gtacatcgcg gtcatcctgg agaacttcag tgttgctact 5400
134 gaagaaagtg cagagcetet gagtgaggat gaetttgaga tgttetatga ggtttgggag 5460
135 aaqtttgatc ccqatqcgac ccaqtttata gagtttgcca aactttctga ttttgcagat 5520
136 gccctggatc ctcctcttct catagcaaaa cccaacaaag tccagctcat tgccatggat 5580
137 ctgcccatgg tgagtggtga ccggatccac tgtcttgaca tcttatttgc ttttacaaag 5640
138 cgtgttttgg gtgagagtgg agagatggat gcccttcgaa tacagatgga agagcgattc 5700
139 atqqcatcaa acccctccaa aqtctcttat qaqcccatta cgaccacqtt gaaacgcaaa 5760
140 caagaggagg tgtctgctat tattatccag agggcttaca gacgctacct cttgaagcaa 5820
141 aaagttaaaa aggtatcaag tatatacaag aaagacaaag gcaaagaatg tgatggaaca 5880
142 cccatcaaag aagatactct cattgataaa ctgaatgaga attcaactcc agagaaaacc 5940
143 gatatgaege ettecaceae gtetecacee tegtatgata gtgtgaecaa accagaaaaa 6000
144 gaaaaatttg aaaaagacaa atcagaaaag gaagacaaag ggaaagatat cagggaaagt 6060
145 aaaaagtaaa aagaaaccaa gaattttcca ttttgtgatc aattgtttac agcccgtgat 6120
146 ggtgatgtgt ttgtgtcaac aggactccca caggaggtct atgccaaact gactgttttt 6180
147 acaaatgtat acttaaggtc agtgcctata acaagacaga gacctctggt cagcaaactg 6240
148 qaactcagta aactggagaa atagtatcga tgggaggttt ctattttcac aaccagctga 6300
149 cactgctgaa gagcagaggc gtaatggcta ctcagacgat aggaaccaat ttaaaggggg 6360
150 gagggaagtt aaatttttat gtaaattcaa catgtgacac ttgataatag taattgtcac 6420
151 cagtgtttat gttttaactg ccacacctgc catattttta caaaacgtgt gctgtgaatt 6480
152 tatcactttt ctttttaatt cacaggttgt ttactattat atgtgactat ttttgtaaat 6540
154 tataactgga tatattttaa atggaggcat gctgcaattc tcattcacac ataaaaaaat 6660
155 cacatcacaa aagggaagag tttacttctt gtttcaggat gtttttagat ttttgaggtg 6720
```

RAW SEQUENCE LISTING DATE: 09/06/2005
PATENT APPLICATION: US/10/501,814 TIME: 11:28:50

```
156 cttaaatagc tattcgtatt tttaaggtgt ctcatccaga aaaaatttaa tgtgcctgta 6780
157 aatgttccat agaatcacaa gcattaaaga gttgttttat ttttacataa cccattaaat 6840
158 gtacatgtat atatgtatat atgtatatgt gcgtgtatat acatatatat gtatacacac 6900
159 atgcacacac agagatatac acataccatt acattgtcat tcacagtccc agcagcatga 6960
161 taagaaqcct gaattgacca aaaaacatcc ccaccaccac tttataaagt tgattctgct 7080
162 ttatcctgca gtattgttta gccatcttct gctcttggta aggttgacat agtatatgtc 7140
163 aatttaaaaa ataaaagtct gctttgtaaa tagtaatttt acccagtggt gcatgtttga 7200
164 gcaaacaaaa atgatgattt aagcacacta cttattgcat caaatatgta ccacagtaag 7260
165 tataqtttqc aaqctttcaa caqqtaatat qatqtaattq qttccattat agtttqaaqc 7320
166 tgtcactgct gcatgtttat cttgcctatg ctgctgtatc ttattccttc cactgttcag 7380
167 aagtctaata tgggaagcca tatatcagtg gtaaagtgaa gcaaattgtt ctaccaagac 7440
168 ctcattcttc atgtcattaa gcaataggtt gcagcaaaca aggaagagct tcttgctttt 7500
169 tattcttcca accttaattg aacactcaat gatgaaaagc ccgactgtac aaacatgttg 7560
170 caagetgett aaatetgttt aaaatatatg gttagagttt tetaagaaaa tataaataet 7620
171 gtaaaaagtt cattttattt tattttcag ccttttgtac gtaaaatgag aaattaaaag 7680
172 tatcttcagg tggatgtcac agtcactatt gttagtttct gttcctagca cttttaaatt 7740
173 gaagcacttc acaaaataag aagcaaggac taggatgcag tgtaggtttc tgctttttta 7800
174 ttagtactgt aaacttgcac acatttcaat gtgaaacaaa tctcaaactg agttcaatgt 7860
175 ttatttgctt tcaatagtaa tgccttatca ttgaaagagg cttaaagaaa aaaaaaatca 7920
176 gctgatactc ttggcattgc ttgaatccaa tgtttccacc tagtcttttt attcagtaat 7980
177 catcagtctt ttccaatgtt tgtttacaca gatagatctt attgacccat atggcactag 8040
178 aactgtatca gatataatat gggatcccag ctttttttcc tctcccacaa aaccaggtag 8100
179 tgaagttata ttaccagtta cagcaaaata ctttgtgttt cacaagcaac aataaatgta 8160
180 gattetttat aetgaageta ttgaettgta gtgtgttggt gaaatgeatg caggaaaatg 8220
181 ctgttaccat aaagaacggt aaaccacatt acaatcaagc caaaagaata aaggtttcgc 8280
182 ttttgttttt gt
                                                                     8292
185 <210> SEQ ID NO: 3
186 <211> LENGTH: 2005
187 <212> TYPE: PRT
188 <213> ORGANISM: Homo sapiens
190 <400> SEQUENCE: 3
191 Met Ala Gln Ser Val Leu Val Pro Pro Gly Pro Asp Ser Phe Arg Phe
                     5
                                        10
192
     1
194 Phe Thr Arg Glu Ser Leu Ala Ala Ile Glu Gln Arg Ile Ala Glu Glu
195
197 Lys Ala Lys Arg Pro Lys Gln Glu Arg Lys Asp Glu Asp Asp Glu Asn
198
            35
200 Gly Pro Lys Pro Asn Ser Asp Leu Glu Ala Gly Lys Ser Leu Pro Phe
203 Ile Tyr Gly Asp Ile Pro Pro Glu Met Val Ser Val Pro Leu Glu Asp
204
                                            75
206 Leu Asp Pro Tyr Tyr Ile Asn Lys Lys Thr Phe Ile Val Leu Asn Lys
207
                                        90
209 Gly Lys Ala Ile Ser Arg Phe Ser Ala Thr Pro Ala Leu Tyr Ile Leu
210
               100
                                   105
                                                       110
212 Thr Pro Phe Asn Pro Ile Arq Lys Leu Ala Ile Lys Ile Leu Val His
                               120
215 Ser Leu Phe Asn Met Leu Ile Met Cys Thr Ile Leu Thr Asn Cys Val
```

RAW SEQUENCE LISTING DATE: 09/06/2005
PATENT APPLICATION: US/10/501,814 TIME: 11:28:50

216	130	)				135					140				
218 Pł	ne Met	Thr	Met	Ser	Asn	Pro	Pro	Asp	Trp	Thr	Lys	Asn	Val	Glu	Tyr
219 14	15				150					155					160
221 Th	ır Phe	Thr	Gly	Ile	Tyr	Thr	Phe	Glu	Ser	Leu	Ile	Lys	Ile	Leu	Ala
222				165					170					175	
224 Aı	rg Gly	Phe	Cys	Leu	Glu	Asp	Phe	Thr	Phe	Leu	Arg	Asp	Pro	Trp	Asn
225			180					185					190		
227 Tı	rp Lei	Asp	Phe	Thr	Val	Iļe	Thr	Phe	Ala	Tyr	Val	Thr	Glu	Phe	Val
228		195					200					205			
230 As	sp Lei	ıGly	Asn	Val	Ser	Ala	Leu	Arg	Thr	Phe	Arg	Val	Leu	Arg	Ala
231	210					215					220				
233 Le	eu Lys	Thr	Ile	Ser	Val	Ile	Pro	Gly	Leu	Lys	Thr	Ile	Val	Gly	Ala
234 22	25				230					235				•	240
236 Le	eu Ile	Gln	Ser	Val	Lys	Lys	Leu	Ser	Asp	Val	Met	Ile	Leu	Thr	Val
237				245					250					255	
239 Pl	ne Cys	Leu	Ser	Val	Phe	Ala	Leu	Ile	Gly	Leu	Gln	Leu	Phe	Met	Gly
240			260			•		265					270		
242 As	sn Lei	ı Arg	Asn	Lys	Cys	Leu	Gln	Trp	Pro	Pro	Asp	Asn	Ser	Ser	Phe
243		275					280					285			_
245 G			Ile	Thr	Ser	Phe	Phe	Asn	Asn	Ser	Leu	Asp	Gly	Asn	Gly
246	290					295					300		_		_
248 Th		Phe	Asn	Arg		Val	Ser	Ile	Phe		Trp	Asp	Glu	Tyr	
249 30		_	_	•	310	_		_		315		_	_		320
251 G	lu As <u>r</u>	Lys	Ser		Phe	Tyr	Phe	Leu		GIY	Gin	Asn	Asp		Leu
252	_	~-3	_	325	_	_		~ 7	330	_	_	<b>~</b> 1	<b>~</b> 3	335	-1.
254 1.4	211 ('VS	1 (2)37	Δcn												
	ou oy.	, сту		ser	ser	Asp	Ата		GIII	Cys	PIO	GIU		Tyr	Ile
255			340					345					350		
255 257 Сչ		Lys	340				Pro	345				Thr	350		
255 257 Cy 258	ys Val	. Lys 355	340 Ala	Gly	Arg	Asn	Pro 360	345 Asn	Tyr	Gly	Tyr	Thr 365	350 Ser	Phe	Asp
255 257 Cy 258 260 Th	ys Val nr Phe	Lys 355 Ser	340 Ala	Gly	Arg	Asn Leu	Pro 360	345 Asn	Tyr	Gly	Tyr Leu	Thr 365	350 Ser	Phe	Asp
255 257 Cy 258 260 Th 261	ys Val nr Phe 370	Lys 355 Ser	340 Ala Trp	Gly Ala	Arg Phe	Asn Leu 375	Pro 360 Ser	345 Asn Leu	Tyr Phe	Gly Arg	Tyr Leu 380	Thr 365 Met	350 Ser Thr	Phe Gln	Asp Asp
255 257 Cy 258 260 Th 261 263 Ph	ys Val nr Phe 370 ne Try	Lys 355 Ser	340 Ala Trp	Gly Ala	Arg Phe Tyr	Asn Leu 375	Pro 360 Ser	345 Asn Leu	Tyr Phe	Gly Arg Arg	Tyr Leu 380	Thr 365 Met	350 Ser Thr	Phe Gln	Asp Asp Thr
255 257 Cy 258 260 Th 261 263 Ph 264 38	ys Val nr Phe 370 ne Try 35	Lys 355 Ser O Glu	340 Ala Trp Asn	Gly Ala Leu	Arg Phe Tyr 390	Asn Leu 375 Gln	Pro 360 Ser Leu	345 Asn Leu Thr	Tyr Phe Leu	Gly Arg Arg 395	Tyr Leu 380 Ala	Thr 365 Met Ala	350 Ser Thr Gly	Phe Gln Lys	Asp Asp Thr 400
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty	ys Val nr Phe 370 ne Try 35	Lys 355 Ser O Glu	340 Ala Trp Asn	Gly Ala Leu Phe	Arg Phe Tyr 390	Asn Leu 375 Gln	Pro 360 Ser Leu	345 Asn Leu Thr	Tyr Phe Leu Phe	Gly Arg Arg 395	Tyr Leu 380 Ala	Thr 365 Met Ala	350 Ser Thr Gly	Phe Gln Lys Tyr	Asp Asp Thr 400
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267	ys Val nr Phe 370 ne Try 35 yr Met	Lys 355 Ser Glu	340 Ala Trp Asn Phe	Gly Ala Leu Phe 405	Arg Phe Tyr 390 Val	Asn Leu 375 Gln Leu	Pro 360 Ser Leu Val	345 Asn Leu Thr	Tyr Phe Leu Phe 410	Gly Arg Arg 395 Leu	Tyr Leu 380 Ala Gly	Thr 365 Met Ala Ser	350 Ser Thr Gly Phe	Phe Gln Lys Tyr 415	Asp Asp Thr 400 Leu
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 269 II	ys Val nr Phe 370 ne Try 35 yr Met	Lys 355 Ser Glu	340 Ala Trp Asn Phe	Gly Ala Leu Phe 405	Arg Phe Tyr 390 Val	Asn Leu 375 Gln Leu	Pro 360 Ser Leu Val	345 Asn Leu Thr	Tyr Phe Leu Phe 410	Gly Arg Arg 395 Leu	Tyr Leu 380 Ala Gly	Thr 365 Met Ala Ser	350 Ser Thr Gly Phe	Phe Gln Lys Tyr 415	Asp Asp Thr 400 Leu
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 269 II	ys Val nr Phe 370 ne Trp 35 yr Met	Lys 355 Ser Glu	340 Ala Trp Asn Phe Ile 420	Gly Ala Leu Phe 405 Leu	Arg Phe Tyr 390 Val Ala	Asn Leu 375 Gln Leu Val	Pro 360 Ser Leu Val	345 Asn Leu Thr Ile Ala 425	Tyr Phe Leu Phe 410 Met	Gly Arg Arg 395 Leu Ala	Tyr Leu 380 Ala Gly Tyr	Thr 365 Met Ala Ser Glu	350 Ser Thr Gly Phe Glu 430	Phe Gln Lys Tyr 415 Gln	Asp Thr 400 Leu Asn
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 269 II	ys Val nr Phe 370 ne Trp 35 yr Met	Lys 355 Ser Glu	340 Ala Trp Asn Phe Ile 420	Gly Ala Leu Phe 405 Leu	Arg Phe Tyr 390 Val Ala	Asn Leu 375 Gln Leu Val	Pro 360 Ser Leu Val	345 Asn Leu Thr Ile Ala 425	Tyr Phe Leu Phe 410 Met	Gly Arg Arg 395 Leu Ala	Tyr Leu 380 Ala Gly Tyr	Thr 365 Met Ala Ser Glu	350 Ser Thr Gly Phe Glu 430	Phe Gln Lys Tyr 415 Gln	Asp Thr 400 Leu Asn
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 269 II 270 272 GJ	ys Val nr Phe 370 ne Try 35 yr Met Le Ass	Lys 355 Ser Glu Ile Leu Thr 435	340 Ala Trp Asn Phe Ile 420 Leu	Gly Ala Leu Phe 405 Leu Glu	Arg Phe Tyr 390 Val Ala Glu	Asn Leu 375 Gln Leu Val	Pro 360 Ser Leu Val Val Glu 440	345 Asn Leu Thr Ile Ala 425 Gln	Tyr Phe Leu Phe 410 Met Lys	Gly Arg Arg 395 Leu Ala Glu	Tyr Leu 380 Ala Gly Tyr	Thr 365 Met Ala Ser Glu Glu 445	350 Ser Thr Gly Phe Glu 430 Phe	Phe Gln Lys Tyr 415 Gln	Asp Thr 400 Leu Asn Gln
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 269 II 270 272 GJ	ys Val nr Phe 370 ne Try 35 yr Met Le Ass	Lys 355 Ser Glu Ile Leu Thr 435	340 Ala Trp Asn Phe Ile 420 Leu	Gly Ala Leu Phe 405 Leu Glu	Arg Phe Tyr 390 Val Ala Glu	Asn Leu 375 Gln Leu Val	Pro 360 Ser Leu Val Val Glu 440	345 Asn Leu Thr Ile Ala 425 Gln	Tyr Phe Leu Phe 410 Met Lys	Gly Arg Arg 395 Leu Ala Glu	Tyr Leu 380 Ala Gly Tyr	Thr 365 Met Ala Ser Glu Glu 445	350 Ser Thr Gly Phe Glu 430 Phe	Phe Gln Lys Tyr 415 Gln	Asp Thr 400 Leu Asn Gln
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 270 272 GJ 273 275 Me 276	ys Val nr Phe 370 ne Trp 35 yr Met Le Ass ln Ala	Lys 355 Ser Glu Ile Leu Thr 435	340 Ala Trp Asn Phe Ile 420 Leu Gln	Gly Ala Leu Phe 405 Leu Glu Leu	Arg Phe Tyr 390 Val Ala Glu Lys	Asn Leu 375 Gln Leu Val Ala Lys 455	Pro 360 Ser Leu Val Val Glu 440 Gln	345 Asn Leu Thr Ile Ala 425 Gln	Tyr Phe Leu Phe 410 Met Lys Glu	Gly Arg 395 Leu Ala Glu Glu	Tyr Leu 380 Ala Gly Tyr Ala Ala 460	Thr 365 Met Ala Ser Glu Glu 445 Gln	350 Ser Thr Gly Phe Glu 430 Phe	Phe Gln Lys Tyr 415 Gln Gln	Asp Thr 400 Leu Asn Gln Ala
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 269 II 270 272 GJ 273 275 Me	ys Val nr Phe 370 ne Trp 35 yr Met le Ass ln Ala 450 la Ala	Lys 355 Ser Glu Ile Leu Thr 435	340 Ala Trp Asn Phe Ile 420 Leu Gln	Gly Ala Leu Phe 405 Leu Glu Leu	Arg Phe Tyr 390 Val Ala Glu Lys	Asn Leu 375 Gln Leu Val Ala Lys 455	Pro 360 Ser Leu Val Val Glu 440 Gln	345 Asn Leu Thr Ile Ala 425 Gln	Tyr Phe Leu Phe 410 Met Lys Glu	Gly Arg 395 Leu Ala Glu Glu	Tyr Leu 380 Ala Gly Tyr Ala Ala 460	Thr 365 Met Ala Ser Glu Glu 445 Gln	350 Ser Thr Gly Phe Glu 430 Phe	Phe Gln Lys Tyr 415 Gln Gln	Asp Thr 400 Leu Asn Gln Ala
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 270 272 Gl 273 275 Me 276 278 Al	ys Val nr Phe 370 ne Trp 35 yr Met Le Ass Ln Ala 450 La Ala	Lys 355 Ser Glu Ile Leu 435 Glu	340 Ala Trp Asn Phe Ile 420 Leu Gln Ser	Gly Ala Leu Phe 405 Leu Glu Leu Ala	Arg Phe Tyr 390 Val Ala Glu Lys Glu 470	Asn Leu 375 Gln Leu Val Ala Lys 455 Ser	Pro 360 Ser Leu Val Val Glu 440 Gln Arg	345 Asn Leu Thr Ile Ala 425 Gln Gln Asp	Tyr Phe Leu Phe 410 Met Lys Glu Phe	Gly Arg 395 Leu Ala Glu Glu Ser 475	Tyr Leu 380 Ala Gly Tyr Ala Ala 460 Gly	Thr 365 Met Ala Ser Glu Glu 445 Gln Ala	350 Ser Thr Gly Phe Glu 430 Phe Ala Gly	Phe Gln Lys Tyr 415 Gln Gln Ala Gly	Asp Asp Thr 400 Leu Asn Gln Ala Ile 480
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 270 272 GJ 273 275 Me 276 278 AJ 279 46	ys Val nr Phe 370 ne Trp 35 yr Met Le Ass Ln Ala 450 La Ala	Lys 355 Ser Glu Ile Leu 435 Glu	340 Ala Trp Asn Phe Ile 420 Leu Gln Ser	Gly Ala Leu Phe 405 Leu Glu Leu Ala	Arg Phe Tyr 390 Val Ala Glu Lys Glu 470	Asn Leu 375 Gln Leu Val Ala Lys 455 Ser	Pro 360 Ser Leu Val Val Glu 440 Gln Arg	345 Asn Leu Thr Ile Ala 425 Gln Gln Asp	Tyr Phe Leu Phe 410 Met Lys Glu Phe	Gly Arg 395 Leu Ala Glu Glu Ser 475	Tyr Leu 380 Ala Gly Tyr Ala Ala 460 Gly	Thr 365 Met Ala Ser Glu Glu 445 Gln Ala	350 Ser Thr Gly Phe Glu 430 Phe Ala Gly	Phe Gln Lys Tyr 415 Gln Gln Ala Gly	Asp Asp Thr 400 Leu Asn Gln Ala Ile 480
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 270 272 GJ 273 275 Me 276 278 AJ 279 46 281 GJ	ys Value 370 ne Try 35 yr Met Le Ass ln Ala et Let 450 la Ala 55	Lys 355 Ser Glu Ile Leu Thr 435 Glu Ala	340 Ala Trp Asn Phe Ile 420 Leu Gln Ser	Gly Ala Leu Phe 405 Leu Glu Leu Ala Glu 485	Arg Phe Tyr 390 Val Ala Glu Lys Glu 470 Ser	Asn Leu 375 Gln Leu Val Ala Lys 455 Ser Ser	Pro 360 Ser Leu Val Val Glu 440 Gln Arg	345 Asn Leu Thr Ile Ala 425 Gln Gln Asp Val	Tyr Phe Leu Phe 410 Met Lys Glu Phe Ala 490	Gly Arg 395 Leu Ala Glu Glu Ser 475 Ser	Tyr Leu 380 Ala Gly Tyr Ala Ala 460 Gly Lys	Thr 365 Met Ala Ser Glu 445 Gln Ala Leu	350 Ser Thr Gly Phe Glu 430 Phe Ala Gly Ser	Phe Gln Lys Tyr 415 Gln Gln Ala Gly Ser 495	Asp Thr 400 Leu Asn Gln Ala Ile 480 Lys
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 270 272 GJ 273 275 Me 276 278 AJ 279 46 281 GJ 282	ys Value 370 ne Try 35 yr Met Le Ass ln Ala et Let 450 la Ala 55	Lys 355 Ser Glu Ile Leu Thr 435 Glu Ala	340 Ala Trp Asn Phe Ile 420 Leu Gln Ser	Gly Ala Leu Phe 405 Leu Glu Leu Ala Glu 485	Arg Phe Tyr 390 Val Ala Glu Lys Glu 470 Ser	Asn Leu 375 Gln Leu Val Ala Lys 455 Ser Ser	Pro 360 Ser Leu Val Val Glu 440 Gln Arg	345 Asn Leu Thr Ile Ala 425 Gln Gln Asp Val	Tyr Phe Leu Phe 410 Met Lys Glu Phe Ala 490	Gly Arg 395 Leu Ala Glu Glu Ser 475 Ser	Tyr Leu 380 Ala Gly Tyr Ala Ala 460 Gly Lys	Thr 365 Met Ala Ser Glu 445 Gln Ala Leu	350 Ser Thr Gly Phe Glu 430 Phe Ala Gly Ser	Phe Gln Lys Tyr 415 Gln Gln Ala Gly Ser 495	Asp Thr 400 Leu Asn Gln Ala Ile 480 Lys
255 257 Cy 258 260 Th 261 263 Ph 264 38 266 Ty 267 270 272 Gl 273 275 Me 276 278 Al 279 46 281 Gl 282 284 Se	ys Value of Phe 370 me Try Met Le Assiln Alaet Leu 450 la Alaet Leu Gluer Glue	Lys 355 Ser Glu Ile Leu Thr 435 Glu Ala Phe	340 Ala Trp Asn Phe 11e 420 Leu Gln Ser Ser Glu 500	Gly Ala Leu Phe 405 Leu Glu Leu Ala Glu 485 Leu	Arg Phe Tyr 390 Val Ala Glu Lys Glu 470 Ser Lys	Asn Leu 375 Gln Leu Val Ala Lys 455 Ser Ser Asn	Pro 360 Ser Leu Val Val Glu 440 Gln Arg Ser Arg	345 Asn Leu Thr Ile Ala 425 Gln Gln Asp Val Arg 505	Tyr Phe Leu Phe 410 Met Lys Glu Phe Ala 490 Lys	Gly Arg 395 Leu Ala Glu Glu Ser 475 Ser Lys	Tyr Leu 380 Ala Gly Tyr Ala Ala 460 Gly Lys Lys	Thr 365 Met Ala Ser Glu 445 Gln Ala Leu Lys	350 Ser Thr Gly Phe Glu 430 Phe Ala Gly Ser Gln 510	Phe Gln Lys Tyr 415 Gln Gln Ala Gly Ser 495 Lys	Asp Thr 400 Leu Asn Gln Ala Ile 480 Lys Glu

VERIFICATION SUMMARY

DATE: 09/06/2005

PATENT APPLICATION: US/10/501,814

TIME: 11:28:51

Input Set : A:\US10501814-seq list.txt Output Set: N:\CRF4\09062005\J501814.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date